

## SEQUENCE LISTING

<110> Andrew, David P.  
Zabel, Brian A.  
Ponath, Paul D.

<120> ANTI-GPR-9-6 ANTIBODIES AND METHODS OF  
IDENTIFYING AGENTS WHICH MODULATE GPR-9-6 FUNCTION

<130> LKS98-16

<140> 09/266,464

<141> 1999-03-11

<160> 7

<170> FastSEQ for Windows Version 3.0

<210> 1

<211> 2577

<212> DNA

<213> Homo sapiens

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<221> CDS

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Ala Asp Asp Tyr Gly Ser Glu Ser Thr Ser Ser Met Glu Asp Tyr Val
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aac ttc aac ttc act gac ttc tac tgt gag aaa aac aat gtc agg cag      156
Asn Phe Asn Phe Thr Asp Phe Tyr Cys Glu Lys Asn Asn Val Arg Gln
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ttt gcg agc cat ttc ctc cca ccc ttg tac tgg ctc gtg ttc atc gtg      204
Phe Ala Ser His Phe Leu Pro Pro Leu Tyr Trp Leu Val Phe Ile Val
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ggg gcc ttg ggc aac agt ctt gtt atc ctt gtc tac tgg tac tgc aca      252
Gly Ala Leu Gly Asn Ser Leu Val Ile Leu Val Tyr Trp Tyr Cys Thr
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aga gtg aag acc atg acc gac atg ttc ctt ttg aat ttg gca att gct      300
Arg Val Lys Thr Met Thr Asp Met Phe Leu Leu Asn Leu Ala Ile Ala
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85 90 95	
gac cag tgg aag ttc cag acc ttc atg tgc aag gtg gtc aac agc atg	396
Asp Gln Trp Lys Phe Gln Thr Phe Met Cys Lys Val Val Asn Ser Met	
100 105 110	
tac aag atg aac ttc tac agc tgt gtg ttg ctg atc atg tgc atc agc	444
Tyr Lys Met Asn Phe Tyr Ser Cys Val Leu Leu Ile Met Cys Ile Ser	
115 120 125	
gtg gac agg tac att gcc att gcc cag gcc atg aga gca cat act tgg	492
Val Asp Arg Tyr Ile Ala Ile Ala Gln Ala Met Arg Ala His Thr Trp	
130 135 140 145	
agg gag aaa agg ctt ttg tac agc aaa atg gtt tgc ttt acc atc tgg	540
Arg Glu Lys Arg Leu Leu Tyr Ser Lys Met Val Cys Phe Thr Ile Trp	
150 155 160	
gta ttg gca gct gct ctc tgc atc cca gaa atc tta tac agc caa atc	588
Val Leu Ala Ala Ala Leu Cys Ile Pro Glu Ile Leu Tyr Ser Gln Ile	
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aag gag gaa tcc ggc att gct atc tgc acc atg gtt tac cct agc gat	636
Lys Glu Glu Ser Gly Ile Ala Ile Cys Thr Met Val Tyr Pro Ser Asp	
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gag agc acc aaa ctg aag tca gct gtc ttg acc ctg aag gtc att ctg	684
Glu Ser Thr Lys Leu Lys Ser Ala Val Leu Thr Leu Lys Val Ile Leu	
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ggg ttc ttc ctt ccc ttc gtg gtc atg gct tgc tgc tat acc atc atc	732
Gly Phe Phe Leu Pro Phe Val Val Met Ala Cys Cys Tyr Thr Ile Ile	
210 215 220 225	
att cac acc ctg ata caa gcc aag aag tct tcc aag cac aaa gcc cta	780
Ile His Thr Leu Ile Gln Ala Lys Lys Ser Ser Lys His Lys Ala Leu	
230 235 240	
aaa gtg acc atc act gtc ctg acc gtc ttt gtc ttg tct cag ttt ccc	828
Lys Val Thr Ile Thr Val Leu Thr Val Phe Val Leu Ser Gln Phe Pro	
245 250 255	
tac aac tgc att ttg ttg gtg cag acc att gac gcc tat gcc atg ttc	876
Tyr Asn Cys Ile Leu Leu Val Gln Thr Ile Asp Ala Tyr Ala Met Phe	
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atc tcc aac tgt gcc gtt tcc acc aac att gac atc tgc ttc cag gtc	924
Ile Ser Asn Cys Ala Val Ser Thr Asn Ile Asp Ile Cys Phe Gln Val	
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acc cag acc atc gcc ttc ttc cac agt tgc ctg aac cct gtt ctc tat	972
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